US ERA ARCHIVE DOCUMENT

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STUDY TITLE

Validation Report for the Determination of Residues of Triclopyr, 3,5,6-Trichloro-2-pyridinol, and 2-Methoxy-3,5,6-trichloropyridine in Fish Tissues by Capillary Gas Chromatography with Mass Selective Detection

DATA REQUIREMENT

171-4(c) & (d)

AUTHORS

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STUDY COMPLETED ON

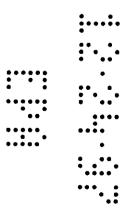
November 24, 1997

PERFORMING LABORATORY

Global Environmental Chemistry Laboratory—Indianapolis Lab
DowElanco
9330 Zionsville Road
Indianapolis, Indiana 46268-1054

LABORATORY STUDY ID

RES94084



STATEMENT OF NO DATA CONFIDENTIALITY CLAIMS

Compound: Triclopyr

Title:	Validation Report for the Determination of Residues of Triclopyr,		
	3,5,6-Trichloro-2-pyridinol, and 2-Methoxy-3,5,6-trichloropyridine in Fish		
	Tissues by Capillary Gas Chromatography with Mass Selective Detection		
	confidentiality is made for any information contained in this study on the basis of it	S	
falling with	n the scope of FIFRA Section 10 (d)(1)(A), (B), or (C).*		
	Company: DowElanco		
	Company Agent: S. A. McMaster		
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STATEMENT OF COMPLIANCE WITH GOOD LABORATORY PRACTICE STANDARDS

Title: Validation Report for the Determination of Residues of Triclopyr, 3,5,6-Trichloro-2pyridinol, and 2-Methoxy-3,5,6-trichloropyridine in Fish Tissues by Capillary Gas Chromatography with Mass Selective Detection

Study Initiation Date: 01-Jul-1994 Study Completion Date: 24-Nov-1997

Experimental Start Date: 01-Jul-1994 Experiment Termination Date: 06-Jul-1997

This report represents data generated after the effective date of the EPA FIFRA Good Laboratory Practice Standards.

United States Environmental Protection Agency Title 40 Code of Federal Regulations Part 160 FEDERAL REGISTER, August 17, 1989

Organisation for Economic Co-Operation and Development ISBN 92-64-12367-9, Paris 1982

All aspects of this study were conducted in accordance with the requirements for Good Laboratory Practice Standards, 40 CFR 160, except as follows: There was no in-progress audit conducted during the study.

S. A. McMaster

Sponsor

DowElanco

S. A. McMaster

Submitter DowElanco

E. L. Olberding

Study Director/Author

DowElanco

Date

Date

November 24 1997

DowElanco Study ID: RES94084 Page 4

DowElanco Quality Assurance Unit Good Laboratory Practice Statement Page

Compound:

Triclopyr

Study ID:

RES94084

Title:

Validation Report for the Determination of Residues of Triclopyr, 3,5,6-Trichloropyr-2-

pyridinol, and 2-Methoxy-3,5,6-trichloropyridine in Fish Tissues by Capillary Gas Cab 11/21/17

Chromatography with Mass Selective Detection

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Study Initiation Date:

01-Jul-1994

Study Completion Date:

11/24/97

24-Nov-1997

GLP Quality Assurance Inspections

Date Reported to Date of GLP the Study Director Inspection(s) and to Management	Phases of the Study which received a GLP Inspection by the Quality Assurance Unit
06-Jun-1994 30-Jun-1994 17, 19-Nov-1997 20-Nov-1997	Protocol Review Report and Raw Data Audit
17, 19-Nov-1997 20-Nov-1997	Report and Raw Data Audi

QUALITY ASSURANCE STATEMENT:

The Quality Assurance Unit has reviewed the final study report and has determined that the report accurately reflects the raw data generated during the conduct of this study.

Thomas G. Lewis

DowElanco, Quality Assurance

Date

SIGNATURE PAGE

Edward & Ollerding	November 24 1997
E. L. Olberding	Date
Study Director/Author	
DowElanco	
DR. Foster Co-Author DowElanco	10/21/97 Date
G. A. Bormett Technical Leader/Reviewer DowElanco	10/23/97 Date
T. E. Weglarz Technical Leader/Reviewer DowElanco	10/27/97 Date
A. S. McGibbon Manager, Analytical Services/Method Development DowElanco	iv 27/97 Date

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Validation Report for the Determination of Residues of Triclopyr, 3,5,6-Trichloro-2-pyridinol, and 2-Methoxy-3,5,6-trichloropyridine in Fish Tissues by Capillary Gas Chromatography with Mass Selective Detection

ABSTRACT

This report contains validation data for DowElanco analytical method GRM 97.02. Residues of triclopyr, 3,5,6-trichloro-2-pyridinol, and 2-methoxy-3,5,6-trichloropyridine were determined in fish tissues by capillary gas chromatography with mass selective detection. The method was validated over the concentration range $0.01-5.0 \,\mu\text{g/g}$ with a limit of quantitation of $0.01 \,\mu\text{g/g}$.

The mean recovery values and standard deviations of the analytes from samples of fish tissues were $93 \pm 6\%$, $92 \pm 9\%$, and $92 \pm 8\%$, for triclopyr, 3,5,6-trichloro-2-pyridinol, and 2-methoxy-3,5,6-trichloropyridine, respectively. The individual recoveries were all within the acceptable range of 70-120% as specified in the study protocol.

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INTRODUCTION

Triclopyr (((3,5,6-trichloro-2-pyridinyl)oxy)acetic acid) is a selective, non-phenoxy herbicide useful in controlling woody plants and broadleaf weeds. Uses of triclopyr-based herbicides include application on rights-of-ways, industrial sites, forest planting sites, turf, range and pasture, and fence rows. The compound is also under development for use as an aquatic herbicide. In the environment, triclopyr degrades primarily to two metabolites, 3,5,6-trichloro-2-pyridinol (3,5,6-TCP), and 2-methoxy-3,5,6-trichloropyridine (2-MP) (1-10). These compounds have also been found in fish and crayfish following exposure to [¹⁴C]triclopyr (11-15).

To support existing and proposed registrations, analytical method GRM 97.02 (Appendix A) has been developed for the determination of residues of triclopyr, 3,5,6-trichloro-2-pyridinol, and 2-methoxy-3,5,6-trichloropyridine in fish tissues by capillary gas chromatography with mass selective detection (GC/MSD) (16). The purpose of the study was to provide validation data to define the accuracy, precision, linearity, and specificity, as well as to support the proposed limit of quantitation. The method was validated over the concentration range of 0.01 to 5.0 μ g/g with a validated method limit of quantitation of 0.01 μ g/g.

EXPERIMENTAL

Sample Origin, Numbering, Preparation, and Storage

Samples of fish tissues were obtained from several sources. The bluegill and catfish samples were collected as part of an aquatic dissipation study (Study Number ENV96052) (17), while the crayfish samples were purchased from Louisiana State University. A portion of these control samples were used for this method validation study.

Upon receipt at DowElanco, the fish tissue samples were inspected and found to be in good condition. Unique sample numbers were assigned to the samples to track them during receipt, storage, and analysis. The fish tissue samples were stored in plastic containers at -15 ° to -20 °C. No significant deviation from this temperature was observed during the storage period between sample receipt and analysis. The sample numbers, sample descriptions, and study numbers are summarized in the table below.

Sample Number	Sample Description	Study Number
SN17993002	bluegill—edible tissue	ENV96052
SN17993003	bluegill—inedible tissue	ENV96052
SN18003302	catfish—edible tissue	ENV96052
SN18003303	catfish—inedible tissue	ENV96052
SN20178201	crayfish—edible tissue	
SN20178202	crayfish—inedible tissue	

Analytical Standards

The test substances and analytical standards used in the validation study were the compounds listed in the following table:

Test Substance/ Analytical Standard	AGR/TSN No.	Percent Purity	Certification Date	Reference
triclopyr ^a	TSN100189	99.52	28-Feb-1997	FA&PC 963272
3,5,6-TCPa	AGO143197	99.97	01-Маг-1995	ML-AL 94-220167
2-MPa	AGR132047	100.0	01-Маг-1995	ML-AL 94-220167
fluroxypyr ^b	AGR222210	99.6	20-Jun-1997	FA&PC 973014
fluroxypyr-DCPb, c	AGR251053	99.5	31-Jul-1995	FA&PC 950258
fluroxypyr-MPb, d	AGR250194	99.9	24-May-1995	FA&PC 950168

a Test substance

Preparation of Triclopyr, 3,5,6-TCP, and 2-MP Spiking Solutions/Calibration Standards

The triclopyr, 3,5,6-TCP, and 2-MP spiking solutions and calibration standards were prepared as described in method GRM 97.02, Section G.1.

Fortification of Recovery Samples

Untreated control samples were fortified with the mixed triclopyr spiking solutions for the generation of method validation recovery data. On the day of analysis, fish tissue samples were fortified by adding $100 \, \mu L$ of the appropriate fortification solution to $1.0 \, g$ of sample. Reagent blank and control samples, containing no mixed triclopyr standard, were also prepared for analysis with the recovery samples to check for background interferences.

b Used as internal standard only

c 4-amino-3,5-dichloro-6-fluoro-2-pyridinol

d 4-amino-3,5-dichloro-6-fluoro-2-methoxypyridine

Sample Extraction and Analysis

Samples were prepared and extracted as described in Section I.1. of analytical method GRM 97.02 with no modifications. Residues of triclopyr, 3,5,6-TCP, and 2-MP were extracted from fish tissues and hydrolyzed using aqueous 0.25 N sodium hydroxide. Following hydrolysis, the sodium hydroxide was acidified and the analytes extracted with ethyl ether. The ethyl ether was passed through an alumina solid-phase extraction (SPE) column which retained the triclopyr and 3,5,6-TCP, while the 2-MP was contained in the ethyl ether eluate. For the determination of 2-MP, the ethyl ether was concentrated and exchanged with 1-chlorobutane. The triclopyr and 3,5,6-TCP were eluted from the alumina column with 0.1 N sodium hydroxide, which was subsequently acidified and purified using a C₁₈ SPE procedure. The eluate from the C₁₈ SPE was then extracted with 1-chlorobutane. For both sample fractions, the 1-chlorobutane was concentrated to less than 1 mL, and an acetone solution containing fluroxypyr analogs as internal standards was added. The samples were then derivatized with *N*-methyl-*N*-(*tert*-butyldimethyl-silyl)-trifluoracetamide (MTBSTFA) to form the *tert*-butyldimethylsilyl (TBDMS) derivatives of triclopyr and 3,5,6-TCP. The samples were then analyzed by capillary GC/MSD.

Calculation of Percent Recovery

Calculation of percent recovery for fortified samples was performed as described in method GRM 97.02, Section I.2.

Statistical Treatment of Data

Statistical treatment of data included the calculation of the least squares regression equations and the correlation coefficients for describing the linearity of the calibration curves, and the calculation of the means, standard deviations, and the relative standard deviations of the results for the fortified recovery samples

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Study Personnel

The following personnel were involved in various aspects of conducting DowElanco Study

Number RES94084: E. L. Olberding, D. R. Foster, M. A. Bartels, D. A. McNett, and

R. D. Griggs.

RESULTS AND DISCUSSION

Radiolabeled Extraction Efficiency Study

For this method validation study, no new extraction efficiency studies were conducted using

radiolabeled analytes. In previous work (11-14), triclopyr, 3,5,6-TCP, 2-MP and their conjugates

were shown to be the major compounds found in fish tissues following exposure to

[14Cltriclopyr. Various extraction solvents and regimens were used in these studies to

completely recover the radioactivity from the samples. In the most recent of these studies (15),

fish tissues heated with aqueous sodium hydroxide for 2 hours at 80 °C were found to extract and

hydrolyze the ¹⁴C-labeled residues. This extraction procedure was adopted in the present

methodology.

Analytical Recovery Data

The method validation study was conducted to determine the recovery levels and the precision of

the method for the determination of residues of triclopyr, 3,5,6-TCP, and 2-MP in fish tissues.

The results are summarized in method GRM 97.02, Tables I-III.

Recovery values of triclopyr from samples of fish tissues fortified over the concentration range of

0.01 to 5.0 µg/g averaged 93% with one standard deviation equal to 6% (Table I). Recovery

values of 3,5,6-TCP from samples of fish tissues fortified over the concentration range of 0.01 to 5.0 µg/g averaged 92% with one standard deviation equal to 9% (Table II). Recovery values of 2-MP from samples of fish tissues fortified over the concentration range of 0.01 to 5.0 µg/g averaged 92% with one standard deviation equal to 8% (Table III). The individual recoveries were all within the acceptable range of 70-120% as specified in the protocol.

Calculated Limits of Quantitation and Detection

Following established guidelines (18), the limits of quantitation (LOQ) and detection (LOD) were calculated using the standard deviation from the 0.01-µg/g recovery results. The LOQ was calculated as ten times the standard deviation (10s), and the LOD was calculated as three times the standard deviation (3s) of the results of the analysis of twelve samples. The results are tabulated in method GRM 97.02, Tables I-III.

The calculated LOQ ranged from 0.007 to 0.009 µg/g for the three analytes, which is lower than the targeted method LOQ of 0.01 µg/g. Results should not be quantified, however, at levels below which no recovery samples have been analyzed.

In a similar fashion, the calculated LOD ranged from 0.002 to 0.003 μ g/g for the three analytes. However, since the lowest level of fortification for recovery samples was 0.003 μ g/g, the method LOD is considered to be 0.003 μ g/g.

Confirmation of Residue Identity

Confirmation of the presence of residues of the analytes is described in method GRM 97.02, Sections I.2.b. and K.2. For the three analytes, confirmation is by comparison of the retention time (gas chromatography) as well as the peak area ratios resulting from selected ion monitoring (mass spectrometry). Confirmation of the presence of the analytes is indicated when the

retention times match those of the standards and the confirmation ratio is in the range of $\pm 20\%$ of the average found for the standards. If additional confirmation is required beyond that discussed in this method, the mass spectra of triclopyr-TBDMS and triclopyr-MP contain additional ions that may be used for confirmation. The confirmation ratios of the samples analyzed during the method validation all fell within $\pm 10\%$ of the average found for the calibration standards.

Representative Full-Scan Mass Spectra

Typical full-scan mass spectra of the TBDMS derivatives of triclopyr and 3,5,6-TCP, and 2-MP are shown in method GRM 97.02, Figures 3-5. Typical full-scan mass spectra of the TBDMS derivatives of fluroxypyr and fluroxypyr-DCP, and fluroxypyr-MP are shown in method GRM 97.02, Figures 3-5.

Representative Calibration Curves

Typical calibration curves for the determination of triclopyr, 3,5,6-TCP, and 2-MP in fish tissues are shown in method GRM 95.19, Figures 6-8.

Representative Chromatograms

Typical chromatograms of a standard, control sample, and a 0.01-µg/g recovery sample for the determination of triclopyr, 3,5,6-TCP, and 2-MP in fish tissues are illustrated in method GRM 97.02, Figures 9-17, respectively. None of the control samples in the method validation study contained interference peaks at the retention times of the analytes or internal standards.

Representative SPE Elution Profiles

Typical elution profiles for 2-MP, triclopyr, and 3,5,6-TCP on an alumina SPE column are illustrated in method GRM 97.02, Figure 18. Typical elution profiles for triclopyr and 3,5,6-TCP on a C₁₈ SPE column are illustrated in method GRM 97.02, Figure 19.

CONCLUSIONS

DowElanco analytical method GRM 97.02, "Determination of Residues of Triclopyr, 3,5,6-Trichloro-2-pyridinol, and 2-Methoxy-3,5,6-trichloropyridine in Fish Tissues by Capillary Gas Chromatography with Mass Selective Detection", has been demonstrated to be suitable for the determination of these analytes in fish tissues. The method was validated over the concentration range of 0.01 to 5.0 μ g/g with a validated method LOQ of 0.01 μ g/g. The individual recoveries for each of the fortified samples were all within the acceptable range of 70-120% as specified in the study protocol.

ARCHIVING

The protocol, raw data, and final report are all filed in the DowElanco testing facility archives at 9330 Zionsville Road, Indianapolis, Indiana 46268-1054.

ACKNOWLEDGEMENTS

Initial sample documentation, handling, and preparation were conducted by R. D. Griggs.

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APPENDIX A

DowElanco Method GRM 97.02

GRM: 97.02

EFFECTIVE: August 19, 1997

SUPERSEDES: ACR 77.4.S1

Determination of Residues of Triclopyr, 3,5,6-Trichloro-2-pyridinol, and 2-Methoxy-3,5,6-trichloropyridine in Fish Tissues by Capillary Gas Chromatography with Mass Selective Detection

E. L. Olberding and D. R. Foster Global Environmental Chemistry Laboratory-Indianapolis Lab DowElanco LLC Indianapolis, Indiana 46268-1054

Scope A.

This method is applicable for the quantitative determination of residues of triclopyr (((3,5,6-trichloro-2-pyridinyl)oxy)acetic acid) and its metabolites, 3,5,6-trichloro-2-pyridinol (3,5,6-TCP), and 2-methoxy-3,5,6-trichloropyridine (2-MP) in fish tissues. The method measures total levels of the analytes (both free and conjugated) over the concentration range 0.01-5.0 µg/g with a validated limit of quantitation of 0.01 µg/g.

B. Principle

Residues of triclopyr, 3,5,6-TCP, and 2-MP are extracted from fish tissues and hydrolyzed using aqueous 0.25 N sodium hydroxide. Following hydrolysis, the sodium hydroxide is acidified and the analytes extracted with ethyl ether. The ethyl ether is passed through an alumina solid-phase extraction (SPE) column which retains the triclopyr and 3,5,6-TCP. while the 2-MP is contained in the ethyl ether eluate. For the determination of 2-MP, the ethyl ether is concentrated and exchanged with 1-chlorobutane. The triclopyr and 3,5,6-TCP are eluted from the alumina column with 0 1 N sodium hydroxide, which is subsequently

acidified and purified using a C₁₈ SPE procedure. The cluate from the C₁₈ SPE is then extracted with 1-chlorobutane. For both sample fractions, the 1-chlorobutane is concentrated to less than 1 mL, and an acetone solution containing fluroxypyr analogs as internal standards is added. The samples are then derivatized with N-methyl-N-(tert-butyldimethylsilyl)-trifluoracetamide (MTBSTFA) to form the tert-butyldimethylsilyl (TBDMS) derivatives of triclopyr and 3,5,6-TCP. The samples are then analyzed by capillary gas chromatography with mass selective detection.

C. Safety Precautions

- Each analyst must be acquainted with the potential hazards of the reagents, products, and solvents used in this method before commencing laboratory work. SOURCES OF INFORMATION INCLUDE: MATERIAL SAFETY DATA SHEETS, LITERATURE, AND OTHER RELATED DATA. Safety information on non-DowElanco products should be obtained from the container label or from the supplier.
- Disposal of reagents, reactants, and solvents must be in compliance with local, state, and federal laws and regulations.
- 2. Acetone, acetonitrile, 1-chlorobutane, ethyl ether, and methanol are flammable and should be used in well-ventilated areas away from ignition sources.
- Hydrochloric acid and sodium hydroxide are corrosive and can cause severe burns. It is imperative that proper eye and personal protection equipment be worn when handling these reagents.

D. Equipment (Note L.1.)

- Balance, analytical, Model AE200, Mettler Instrument Corporation, Hightstown, NJ 08520.
- 2. Balance, pan, Model BB2440, Mettler Instrument Corporation.
- 3. Bath, water, catalog number 15-461-20, Fisher Scientific, Pittsburgh, PA 15219.
- 4. Centrifuge, with rotor to accommodate 40-mL vials, Model Centra-8, International Equipment Company, Needham Heights, MA 02194.
- Evaporator, N-Evap, Model 111, Organomation Associates, Inc., South Berlin, MA 01549. (Note L.2.)
- 6. Gas chromatograph, Model 5890A Series II, Hewlett-Packard, Wilmington, DE 19808.
- 7. Hammer mill, with 3/16-inch screen, Model 2001, AGVISE Laboratories, Inc., Northwood, ND 58267.
- 8. Injector, automatic, Model 7673, Hewlett-Packard.

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Effective Date: August 19, 1997 GRM 97.02

9. Mass selective detector, Model 5972A, Hewlett-Packard, Palo Alto, CA 94304.

- 10. Mass selective detector data system, Model G1034C, Hewlett-Packard.
- 11. Oven, Model OV-490A-2, Blue M Electric Company, Blue Island, IL 60406.
- 12. Shaker, variable speed reciprocating with box carrier, Model 6000, Eberbach Corporation, Ann Arbor, MI 48106.
- Ultrasonic cleaner, Model 1200, Branson Cleaning Equipment Company, Shelton, CT 06484.
- 14. Vacuum manifold, Model spe-12G, Mallinckrodt Baker, Inc., Phillipsburg, NJ 08865.
- 15. Vortex mixer, Model G-560, Scientific Industries, Inc., Bohemia, NY 11716.
- Water purification system, Model Milli-Q UV Plus, Millipore Corporation, Milford, MA 01757.

E. Glassware and Materials (Note L.1.)

- Column, capillary gas chromatography, Durabond-1701 liquid phase, 10 m x 0.18 mm i.d., 0.4-μm film thickness, catalog number 121-0713, J & W Scientific, Folsom, CA 95630.
- 2. Column, alumina SPE, catalog number 7214-07, Mallinckrodt Baker, Inc.
- 3. Column, C₁₈ SPE, catalog number 7020-07, Mallinckrodt Baker, Inc.
- 4. Cylinder, graduated mixing, 1000-mL, catalog number 6224-27, Ace Glass Inc., Vineland, NJ 08360.
- 5. Filter, charcoal, catalog number 7972, Chrompack, Inc., Raritan, NJ 08869. (Note L.3.)
- 6. Filter, moisture, catalog number 7971, Chrompack, Inc. (Note L.3.)
- 7. Filter, oxygen, catalog number 7970, Chrompack, Inc. (Note L.3.)
- 8. Flask, volumetric, 100-mL, catalog number 161-8987, National Scientific Company, Lawrenceville, GA 30243.
- 9. Flask, volumetric, 200-mL, catalog number 161-8988, National Scientific Company.
- 10. Inlet sleeve, double gooseneck splitless, catalog number 5181-3315, Hewlett-Packard.
- 11. pH indicator sticks, pH range 0-6, catalog number 4391-01, Mallinckrodt Baker, Inc.

- 12. Pipet, serological, 1.0-mL, catalog number P4760-1C, National Scientific Company.
- 13. Pipet, volumetric, 1.0-mL, catalog number 261-6011, National Scientific Company.
- 14. Pipet, volumetric, 2.0-mL, catalog number 261-6012, National Scientific Company.
- 15. Pipet, volumetric, 2.5-mL, catalog number 261-6084, National Scientific Company.
- 16. Pipet, volumetric, 3.0-mL, catalog number 261-6013, National Scientific Company.
- 17. Pipet, volumetric, 4.0-mL, catalog number 261-6014, National Scientific Company.
- 18. Pipet, volumetric, 5.0-mL, catalog number 261-6015, National Scientific Company.
- 19. Pipet, volumetric, 10-mL, catalog number 261-6020, National Scientific Company.
- 20. Pipet, volumetric, 20-mL, catalog number 261-6030, National Scientific Company.
- 21. Pipet, volumetric, 25-mL, catalog number 261-6035, National Scientific Company.
- 22. Pipet, volumetric, 50-mL, catalog number 261-6050, National Scientific Company.
- 23. Syringe, 50-μL, Model 705N, Hamilton Company, Reno, NV 89520.
- 24. Syringe, 100-μL, Model 710N, Hamilton Company.
- 25. Syringe, 250-µL, Model 725N, Hamilton Company.
- 26. Syringe, 500-µL, Model 750N, Hamilton Company.
- 27. Vial, autosampler, 2-mL, catalog number C4000-1, National Scientific Company.
- 28. Vial, 12-mL, with PTFE-lined screw cap, catalog number B7800-12, National Scientific Company.
- 29. Vial, 16-mL, with PTFE-lined screw cap, catalog number B7800-4, National Scientific Company.
- 30. Vial, 40-mL, with PTFE-lined screw cap, catalog number B7800-6, National Scientific Company.
- 31. Vial cap, for autosampler vial, catalog number C4000-54B, National Scientific Company.

F. Reagents and Chemicals (Note L.1.)

1. Reagents

- Acetone, OmniSolv grade, catalog number AX0110-1, EM Science, Gibbstown, NJ 08027.
- b. Acetonitrile, OmniSolv grade, catalog number AX0151-1, EM Science.
- c. 1-Chlorobutane, OmniSolv grade, catalog number CX0914-1, EM Science.
- d Ethyl ether, anhydrous, catalog number EX0190-6, EM Science.
- e. Helium, gas, 99.995% purity, Airco, Murray Hıll, NJ 07974.
- f. Hydrochloric acid, 5.0 N, catalog number LC15360-2, Fisher Scientific.
 - g. Hydrochloric acid, 1.0 N, catalog number SA48-1, Fisher Scientific.
 - h. Hydrochloric acid, 0.10 N, catalog number SA54-1, Fisher Scientific.
 - i. Methanol, OmniSolv grade, catalog number MX-0480-1, EM Science.
 - j. MTBSTFA (*N*-methyl-*N*-(*tert*-butyldimethylsilyl)-trifluoroacetamide), catalog number 48920, Pierce Chemical Company, Rockford, IL 61105.
 - k. Nitrogen, gas, 99.99% purity, Airco.
 - Nitrogen, refrigerated liquid, catalog number LQNI-230, Airco Gas and Gear, Indianapolis, IN 46241.
 - m. Sodium chloride, ACS reagent grade, catalog number S271-1, Fisher Scientific.
 - n. Sodium hydroxide, 0.25 N, certified concentration, catalog number SS272-1, Fisher Scientific.
 - Sodium hydroxide, 0.10 N, certified concentration, catalog number SS276-1, Fisher Scientific.

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p. Standards

- (1) triclopyr (((3,5,6-trichloro-2-pyridinyl)oxy)acetic acid)
- (2) 3,5,6-trichloro-2-pyridinol (3,5,6-TCP)
- (3) 2-methoxy-3,5,6-trichloropyridine (2-MP)
- (4) fluroxypyr (((4-amino-3,5,-dichloro-6-fluoro-2-pyridinyl)oxy)acetic acid)
- (5) 4-amino-3,5-dichloro-6-fluoro-2-pyridinol (fluroxypyr-DCP)
- (6) 4-amino-3,5-dichloro-6-fluoro-2-methoxypyridine (fluroxypyr-MP)

Obtain from Test Substance Coordinator, DowElanco, 9330 Zionsville Road, Building 306/A1, Indianapolis, IN 46268-1054.

2. Prepared Solutions

a. 80% acetonitrile/19% water/1% 1.0 N hydrochloric acid solution (v/v/v).

Pour 800 mL of acetonitrile into a 1000-mL graduated mixing cylinder. Pipet 10.0 mL of 1.0 N hydrochloric acid into the same cylinder; then add approximately 150 mL of water. Swirl the cylinder, and allow to equilibrate to room temperature. Dilute to volume with water.

b. 40% acetonitrile/59% water/1% 1.0 N hydrochloric acid solution (v/v/v).

Pour 400 mL of acetonitrile into a 1000-mL graduated mixing cylinder. Pipet 10.0 mL of 1.0 N hydrochloric acid into the same cylinder; then add approximately 500 mL of water. Swirl the cylinder, and allow to equilibrate to room temperature. Dilute to volume with water.

G. Preparation of Standards

1. Preparation of Spiking Solutions/Calibration Standards

- Weigh 0.1000 g of triclopyr analytical standard and quantitatively transfer to a 100-mL volumetric flask. Dilute to volume with acetone to obtain a 1000-μg/mL stock solution.
- b. Weigh 0.1000 g of 3,5,6-trichloro-2-pyridinol analytical standard and quantitatively transfer to a 100-mL volumetric flask. Dilute to volume with acetone to obtain a 1000-μg/mL stock solution.
- c. Weigh 0.1000 g of 2-methoxy-3,5,6-trichloropyridine analytical standard and quantitatively transfer to a 100-mL volumetric flask. Dilute to volume with acetone to obtain a 1000-μg/mL stock solution.

d. Pipet 20.0 mL of each of the stock solutions in Sections G.1.a.-c. into a single 200-mL volumetric flask and adjust to volume with acetone to obtain a solution containing 100.0 μg/mL of each compound

e. Prepare solutions for spiking fish tissues by diluting the solution from Section G.1.d with acetone as follows:

Aliquot of Initial Soln.	Final Soln. Volume	Spiking Soln. Final Conc.	Equivalent Sample Conc. ^a
mL	mL	μg/mL	μg/g
0.031	100	0 031	0.003
0.050	100	0.050	0.005
0.100	100	0.100	0.010
0.250	100	0.250	0.025
0.500	100	0.500	0.050
1.00	100	1.00	0.100
2.50	100	2.50	0.250
5.00	100	5.00	0.500
10.0	100	10.0	1.00
25.0	100	25.0	2.50
50.0	100	50.0	5.00

The equivalent sample concentration is based on fortifying a 1.0-g fish tissue sample with 100 μL of spiking solution.

f. Prepare calibration standards by dispensing 100 μL of the solutions from Section G.1.e. into 12-mL vials containing 0.5 mL of 1-chlorobutane and derivatizing according to the procedure described in Section I.1.ii.-mm. The concentration range of these calibration standards is from 0.005 to 5.0 μg/mL.

Chemical structures of the underivatized and derivatized triclopyr, 3,5,6-TCP, and 2-MP are shown in Figure 1.

2. Preparation of Internal Standard Solution

- a. Weigh 0.1000 g of fluroxypyr analytical standard and quantitatively transfer to a 100-mL volumetric flask. Dilute to volume with acetone to obtain a 1000-μg/mL stock solution.
- b. Weigh 0.1000 g of 4-amino-3,5-dichloro-6-fluoro-2-pyridinol analytical standard and quantitatively transfer to a 100-mL volumetric flask. Dilute to volume with acetone to obtain a 1000-μg/mL stock solution.
- c. Weigh 0.1000 g of 4-amino-3,5-dichloro-6-fluoro-2-methoxypyridine analytical standard and quantitatively transfer to a 100-mL volumetric flask. Dilute to volume with acetone to obtain a 1000-µg/mL stock solution.

d. Pipet 20.0 mL of each of the stock solutions in Sections G.2.a.-c. into a single 200-mL volumetric flask and adjust to volume with acetone to obtain a solution containing 100.0 μg/mL of each compound.

e. Pipet 20.0 mL of the solution in Section G.2.d. into a 200-mL volumetric flask and adjust to volume with acetone to obtain a mixture containing 10.0 μg/mL of each compound

Chemical structures of the underivatized and derivatized fluroxypyr, fluroxypyr-DCP, and fluroxypyr-MP are shown in Figure 2.

H. Gas Chromatography/Mass Spectrometry

1. Column

Install the splitless column inlet sleeve (Section E.10.) and the capillary column (Section E.1.) in the split/splitless injection port of the gas chromatograph following the manufacturer's recommended procedures.

2. Typical Operating Conditions (Note L.4.)

Instrumentation: Hewlett-Packard Model 5890A Gas Chromatograph

Hewlett-Packard Model 7673 Automatic Sampler Hewlett-Packard Model 5972A Mass Selective Detector Hewlett-Packard Model G1034C Data System Software

Column: J & W Scientific fused silica capillary

Durabond-1701 liquid phase

10 m x 0.18 mm i.d. 0.4-µm film thickness

Temperatures:

Column 70 °C for 1.0 min

70 °C to 255 °C at 10 °C/min 255 °C to 280 °C at 20 °C/min

280 °C for 4.75 min

Injector 280 °C Interface 280 °C

Carrier Gas: helium

Constant Flow on Vacuum Compensation on Head Pressure 25 kPa

Linear Velocity approximately 45 cm/s

Injection Mode: splitless

Purge Delay 0.9 min
Splitter Flow 50 mL/min
Septum Purge 1.0 mL/min

Injection Volume: $1 \mu L$

Detector: electron impact selected ion monitoring

Calibration Program maximum sensitivity autotune (Note L.5.)
Electron Multiplier 2050 volts (≈280 volts above autotune)

Ions Monitored: (Note L.6.)

Triclopyr-TBDMS m/z 314 (quantitation)

m/z 254, 256, 312 (confirmation) (Section K.2.)

3,5,6-TCP-TBDMS m/z 254 (quantitation)

m/z 256 (confirmation)

2-MP m/z 213 (quantitation)

m/z 182, 210, 211, 212 (confirmation) (Section K.2.)

Fluroxypyr-TBDMS m/z 311 (internal standard for triclopyr-TBDMS)
Fluroxypyr-DCP-TBDMS m/z 253 (internal standard for 3,5,6-TCP-TBDMS)

Fluroxypyr-MP m/z 210 (internal standard for 2-MP)

Dwell Time 75 ms

Mass spectra of the above triclopyr and fluroxypyr compounds are shown in Figures 3-5, respectively.

3. Calibration Curves

Typical calibration curves for the determination of triclopyr, 3,5,6-TCP, and 2-MP in fish tissues are shown in Figures 6-8, respectively.

4. Typical Chromatograms

Typical chromatograms of a standard, control sample, and a 0.01-µg/g recovery sample for the determination of triclopyr, 3,5,6-TCP, and 2-MP in fish tissues are illustrated in Figures 9-17, respectively. Crayfish inedible tissue was chosen since it represents the "worst case" situation for sample preparation and analysis.

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I. <u>Determination of Recovery of Triclopyr and Metabolites from Fish Tissues</u>

1. Preparation of Recovery Samples

- a. Prepare the tissue sample for analysis by freezing with liquid nitrogen and then grinding the sample using a hammer mill with a 3/16-inch screen.
- b. Accurately weigh 1.0-g portions of the prepared fish tissue sample into a series of 40-mL vials.
- c. For preparing fortified samples, use some of the samples as controls and fortify the remaining samples by adding 100-μL aliquots of the appropriate spiking solutions (Section G.1.e.) to obtain concentrations ranging from 0.01 to 5.0 μg/g. A reagent blank, containing no fish tissue, should be carried through the method with the samples.
- d. Add 10.0 mL of the 0.25 N sodium hydroxide extraction solution to the sample vial.
- e. Cap the sample vial with a PTFE-lined cap and vortex the sample for 5-10 seconds.
- f. Hydrolyze the sample by heating in a water bath for 2 hours at 80 °C.
- g. After hydrolysis, allow the sample to cool to room temperature.
- h. Using a serological pipet, add 0.75 mL of 5.0 N hydrochloric acid to the sample vial and swirl gently to mix. Measure the pH to insure that the pH<1. If necessary, adjust the pH by the dropwise addition of 5.0 N hydrochloric acid. (Note L.7.)
- i. Add 5 g of sodium chloride (enough to saturate the solution) and 10 mL of ethyl ether to the sample vial.
- j. Cap the sample vial with a PTFE-lined cap, and shake the sample for a minimum of 1 hour on a reciprocating shaker at approximately 250 excursions/minute.
- k. Centrifuge the sample vial for 5 minutes at approximately 2000 rpm.
- 1. Transfer the ethyl ether (top) layer into a clean 16-mL vial.
- m. Add an additional 5 mL of ethyl ether to the sample vial. Cap the vial, and shake the sample for an additional hour on a reciprocating shaker at approximately 250 excursions/minute.
- n. Centrifuge the sample vial for 5 minutes at approximately 2000 rpm.

o. Transfer the ethyl ether layer from Step I.1.n. into the same vial containing the ethyl ether from Step I.1.l.

- p. Repeat Steps I.1.m.-o. one additional time. After transferring the ethyl ether layer into the 16-mL vial from the first extraction, discard the 40-mL vial containing the aqueous portion of the sample extract.
- q. Concentrate the solution from Step I.1.p. to 5 ± 0.5 mL using an N-Evap evaporator. (Note L.8.)
- r. Purify the sample using the following alumina SPE procedure (Sections K.4.a. and K 4.b.):
- (1) Place an alumina SPE column on the vacuum manifold.
- (2) Rinse the SPE column with 5 mL of methanol. (Do not allow the column bed to dry.)
- (3) Condition the SPE column with 5 mL of ethyl ether. (Do not allow the column bed to dry.)
- (4) Place a 12-mL vial in the vacuum manifold box to collect the eluate in the following step.
- (5) Transfer the sample solution from Step I.1.q. to the SPE column and allow the sample to flow through the column by gravity alone, collecting the eluate in the 12-mL vial. (Do not allow the column bed to dry.)
- (6) Rinse the sample vial with 2 mL of ethyl ether. When the sample solution in Step I.1.r.(5) is within 2 mm of the top of the column bed, transfer the rinse to the SPE column. Allow the solvent to flow through the column by gravity alone, collecting the cluate in the same 12-mL vial.
- (7) Remove the 12-mL vial from the vacuum manifold box and set it aside. (This extract, containing the 2-methoxy-3,5,6-trichloropyridine, will be further treated as described in Steps I.1.s.-x.)
- (8) Rinse the sample vial with 5 mL of acetone and transfer the rinse to the SPE column. Allow the solvent to flow through the column by gravity alone, and discard the eluate.
- (9) Dry the SPE column under vacuum for 5 minutes.
- (10) Place a 40-mL vial in the vacuum manifold box to collect the eluate in the following step.
- (11) Elute the triclopyr and 3,5,6-trichloro-2-pyridinol from the alumina SPE column by passing 25 mL of 0.1 N sodium hydroxide solution through the column at a flow rate of 1-2 mL/min, collecting the eluate in the 40-mL vial. (This extract, containing the triclopyr and 3,5,6-trichloro-2-pyridinol, will be further treated as described in Steps I 1.y.-nn.)

2-Methoxy-3,5,6-trichloropyridine

- s. Concentrate the solution from Step I.1.r.(7) to approximately 2 mL using an N-Evap evaporator. (Note L.8.)
- t. Add 2.0 mL of 1-chlorobutane and continue concentrating the solution from Step I.1.s. to less than 0.8 mL (but not to dryness) using an N-Evap evaporator. (Notes L.2. and L.8.)
- u. Add 100 μ L of the internal standard solution (Section G.2.e) and 100 μ L of MTBSTFA derivatizing reagent to the sample vial.
- v. Adjust the volume in the sample vial to 1.0 mL with 1-chlorobutane and firmly seal with a PTFE-lined cap (Note L.8.). Vortex the sample for 5-10 seconds, and then sonicate the sample for 5-10 seconds.
- w. Transfer the solution to a 2-mL autosampler vial and seal with a cap.
- x. Analyze the calibration standards (Section G.1.f.) and samples by capillary gas chromatography/mass spectrometry as described in Section H.2. Determine the suitability of the chromatographic system using the following performance criteria:
- (1) Standard curve linearity: Determine that the correlation coefficient equals or exceeds 0.995 for the least squares equation which describes the detector response as a function of standard curve concentration. If power regression is used, the power exponent should be between 0.90-1.10.
- (2) Peak resolution: Visually determine that sufficient resolution has been achieved for the analyte and internal standard relative to background interferences.
- (3) Appearance of chromatograms: Visually determine that the chromatograms resemble those shown in Figures 15-17 with respect to peak response, baseline noise, and background interference. Visually determine that a minimum signal-to-noise ratio of 10:1 has been attained for the 2-methoxy-3,5,6-trichloropyridine in the 0.010-µg/mL calibration standard.

Triclopyr and 3,5,6-Trichloro-2-pyridinol

- y. Using a serological pipet, add 1.1 mL of 5.0 N hydrochloric acid to the sample vial from Step I.1.r.(11), cap the vial, and vortex the sample for 5-10 seconds.
- z. Purify the sample using the following C₁₈ SPE (Section K.4.c.):
- (1) Place a C₁₈ SPE column on the vacuum manifold.
- (2) Rinse the SPE column with 5 mL of acetonitrile. (Do not allow the column bed to dry.)

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(3) Condition the SPE column with 5 mL of 0.1 N hydrochloric acid. (Do not allow the column bed to dry.)

- (4) Transfer the sample solution from Step I.1.y. to the SPE column, and slowly pull the sample through the column at a flow rate of 1-2 mL/min with the aid of vacuum. Discard the eluate without allowing the column bed to dry.
- (5) Rinse the sample vial with 4.0 mL of a 40% acetonitrile/59% water/1% 1.0 N hydrochloric acid solution. When the sample solution in Step I.1.z.(4) is within 2 mm of the top of the column bed, transfer the rinse to the SPE column. With the aid of vacuum, slowly pull the rinse solution through the column, and discard the eluate.
- (6) Dry the SPE column under vacuum for 1 minute.
- (7) Place a 40-mL vial in the vacuum manifold box to collect the eluate in the following step.
- (8) Elute the triclopyr and 3,5,6-trichloro-2-pyridinol from the C₁₈ SPE column by passing 3.0 mL of an 80% acetonitrile/19% water/1% 1.0 N hydrochloric acid solution through the column, collecting the eluate in the 40-mL vial.
- aa. Add 10 mL of 0.1 N hydrochloric acid, 5 g of sodium chloride (enough to saturate the solution), and 5 mL of 1-chlorobutane to the sample vial.
- bb. Cap the sample vial with a PTFE-lined cap, and shake the sample for a minimum of 10 minutes on a reciprocating shaker at approximately 180 excursions/minute.
- cc. Centrifuge the sample vial for 5 minutes at approximately 2000 rpm.
- dd Transfer the 1-chlorobutane layer into a clean 12-mL vial. (Note L.9.)
- ee. Add an additional 5 mL of 1-chlorobutane to the sample vial. Cap the vial, and shake the sample for an additional 20 minutes on a reciprocating shaker at approximately 180 excursions/minute.
- ff. Centrifuge the sample vial for 5 minutes at approximately 2000 rpm.
- gg. Transfer the 1-chlorobutane layer from Step I.1.ff. into the same vial containing the 1-chlorobutane from Step I.1.dd. (Note L.9.)
- hh. Concentrate the solution from Step I.1.gg. to less than 0.8 mL (but not to dryness) using an N-Evap evaporator. (Notes L.2. and L.8.)
- 11 Add 100 μ L of the internal standard solution (Section G.2.e) and 100 μ L of MTBSTFA derivatizing reagent to the sample vial.

- jj. Adjust the volume in the sample vial to 1.0 mL with 1-chlorobutane and firmly seal with a PTFE-lined cap (Note L.8.). Vortex the sample for 5-10 seconds, and then sonicate the sample for 5-10 seconds.
- kk. Place the sample vial in an oven at 60 °C and allow the mixture to react for 60 minutes.
- ll. Remove the sample vial from the oven and allow the reaction mixture to cool to room temperature.
- mm. Transfer the solution to a 2-mL autosampler vial and seal with a cap.
- nn. Analyze the calibration standards (Section G.1.f.) and samples by capillary gas chromatography/mass spectrometry as described in Section H.2. Determine the suitability of the chromatographic system using the following performance criteria:
 - (1) Standard curve linearity: Determine that the correlation coefficients equal or exceed 0.995 for the least squares equations which describes the detector response as a function of standard curve concentration. If power regression is used, the power exponents should be between 0.90-1.10.
 - (2) Peak resolution: Visually determine that sufficient resolution has been achieved for the analytes and internal standards relative to background interferences.
 - (3) Appearance of chromatograms: Visually determine that the chromatograms resemble those shown in Figures 9-14 with respect to peak response, baseline noise, and background interference. Visually determine that a minimum signal-to-noise ratio of 10:1 has been attained for each analyte in the 0.010-µg/mL calibration standard.

2. Calculation of Percent Recovery

a. Inject the series of calibration standards described in Section G.1.f. and determine the peak areas for the analytes and internal standards as indicated below.

Triclopyr-TBDMS

3,5,6-TCP-TBDMS

2-MP

m/z 214 (quantitation), m/z 256 (confirmation)

m/z 254 (quantitation), m/z 256 (confirmation)

m/z 213 (quantitation), m/z 211 (confirmation)

m/z 311 (internal standard for triclopyr-TBDMS)

Fluroxypyr-DCP-TBDMS

m/z 253 (internal standard for 3,5,6-TCP-TBDMS)

Fluroxypyr MP

m/z 210 (internal standard for 2-MP)

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b. For each standard, calculate each analyte's confirmation ratio. Use the average confirmation ratio for each analyte to confirm the presence of the analyte in the fish tissue samples.

For example, using the data for triclopyr from Figure 9:

Confirmation Ratio = $\frac{\text{peak area of confirmation ion}}{\text{peak area of quantitation ion}}$

Confirmation Ratio = $\frac{\text{peak area at } m/z \ 312}{\text{peak area at } m/z \ 314}$

Confirmation Ratio = $\frac{16597}{16562}$

Confirmation Ratio = 1.0021

Confirmation of the presence of the analyte is indicated when the confirmation ratio for the sample is within the range of $\pm 20\%$ of the average found for the standards.

c. For each standard, calculate each analyte's quantitation ratio.

For example, using the data for triclopyr from Figure 9:

Quantitation Ratio = $\frac{\text{peak area of quantitation ion}}{\text{peak area of internal standard ion}}$

Quantitation Ratio = $\frac{\text{peak area at } m/z \text{ } 314}{\text{peak area at } m/z \text{ } 311}$

Quantitation Ratio = $\frac{16562}{1723923}$

Quantitation Ratio = 0.00961

d. Prepare a standard curve for each analyte by plotting the equivalent analyte concentration on the abscissa (x-axis) and the respective quantitation ratio on the ordinate (y-axis) as shown in Figures 6-8. Using regression analysis, determine the equation for the curve with respect to the abscissa.

For example, using power regression (1) with the triclopyr data from Figure 6:

$$Y = constant \times X^{(exponent)}$$

$$X = \left(\frac{Y}{constant}\right)^{1/exponent}$$

$$Triclopyr Conc. = \left(\frac{triclopyr quantitation ratio}{constant}\right)^{1/exponent}$$

$$Triclopyr Conc. = \left(\frac{triclopyr quantitation ratio}{0.8925}\right)^{1/1.0139}$$

e. Determine the gross concentration in each recovery sample by substituting the quantitation ratio obtained into the above equation and solving for the concentration.

For example, using the triclopyr data from Figure 11:

Triclopyr Conc. (gross
$$\mu g/g$$
) = $\left(\frac{\text{triclopyr quantitation ratio}}{0.8925}\right)^{1/1.0139}$

Triclopyr Conc. (gross $\mu g/g$) = $\left(\frac{0.0085}{0.8925}\right)^{1/1.0139}$

Triclopyr Conc. = $0.0102 \, \mu g/g$

f. Determine the net concentration in each recovery sample by subtracting the triclopyr concentration in the control sample from that of the gross triclopyr concentration in the recovery sample.

For example, using the triclopyr data from Figures 10 and 11:

Triclopyr Conc.
$$=$$
 Triclopyr Conc. $-$ Triclopyr Conc. $(\text{gross } \mu g/g)$ $(\text{control } \mu g/g)$

Triclopyr Conc. $=$ 0.0102 $\mu g/g$ $-$ 0.0000 $\mu g/g$

Triclopyr Conc. $=$ 0.0102 $\mu g/g$

(net)

g. Determine the percent recovery by dividing the net concentration of each recovery sample by the theoretical concentration added.

Recovery =
$$\frac{\text{Concentration Found}}{\text{Concentration Added}} \times 100\%$$

Recovery =
$$\frac{0.0102 \,\mu g/g}{0.0100 \,\mu g/g} \times 100\%$$

Recovery
$$= 102\%$$

- J. Determination of Triclopyr and Metabolites in Fish Tissues
 - 1. Prepare reagent blank, control, recovery, and treated samples as described in Section I.1.
 - 2. Prepare a standard calibration curve for triclopyr, 3,5,6-TCP, and 2-MP and determine the percent recovery for each analyte as described in Section I.2.
 - 3. Determine the gross concentration of each analyte in each treated sample by substituting the quantitation ratio obtained into the equation for the standard calibration curve, and calculating the uncorrected residue result as described in Section I.2.e.
 - 4. For those analyses that require correction for method recovery, use the average recovery of all the recovery samples from a given sample set to correct for method efficiency.

For example, using the triclopyr data from Figure 11 and Table I for the samples analyzed on 3-Jul-1997:

- Determine the gross analyte concentrations in the fish tissue sample as described in Section I.2.e.
- b. Determine the corrected analyte concentration in the fish tissue sample as follows:

Triclopyr Conc.
$$=$$
 Triclopyr Conc. $\times \left(\frac{100}{\text{Average Percent Recovery}}\right)$

Triclopyr Conc.
$$= 0.0102 \mu g/g \times \frac{100}{92}$$

K. Results and Discussion

1. Method Validation

a. Recovery Levels and Precision

A method validation study was conducted to determine the recovery levels and the precision of the method for the determination of triclopyr, 3,5,6-TCP, and 2-MP in fish tissues. The results are summarized in Tables I-III.

Recovery values of triclopyr from samples of fish tissues fortified over the concentration range of 0.01 to 5.0 μ g/g averaged 93% with one standard deviation equal to 6% (Table I). Recovery values of 3,5,6-TCP from samples of fish tissues fortified over the concentration range of 0.01 to 5.0 μ g/g averaged 92% with one standard deviation equal to 9% (Table II). Recovery values of 2-MP from samples of fish tissues fortified over the concentration range of 0.01 to 5.0 μ g/g averaged 92% with one standard deviation equal to 8% (Table III).

b. Standard Curve Linearity

For the power least squares regression equations describing the detector response as a function of the standard calibration curve concentrations, correlation coefficients (r²) were greater than 0.999 for all three analytes, while the power exponents were between 0.97 and 1.01.

c. Calculated Limits of Quantitation and Detection

Following established guidelines (2), the limits of quantitation (LOQ) and detection (LOD) were calculated using the standard deviation from the 0.01- μ g/g recovery results. The LOQ was calculated as ten times the standard deviation (10s), and the LOD was calculated as three times the standard deviation (3s) of the results of the analysis of 12 samples. The results are summarized in Tables I-III.

For all three analytes, the calculated LOQ ranged from 0.007 to 0.009 $\mu g/g$, which is lower than the targeted method LOQ of 0.01 $\mu g/g$. In a similar fashion, for all three analytes the calculated LOD ranged from 0.002 to 0.003 $\mu g/g$. However, since the lowest level of fortification for recovery samples was 0 003 $\mu g/g$, the method LOD is considered to be 0.003 $\mu g/g$.

2. Confirmation of Residue Identity

Confirmation of the presence of residues is described in Section I.2.b. For the three analytes, confirmation is by comparison of the retention time (gas chromatography) as well as the peak area ratios resulting from selected ion monitoring (mass spectrometry). Confirmation of the presence of the analytes is indicated when the confirmation ratio for the sample is in the range of ±20% of the average found for the standards. If additional confirmation is required beyond that discussed in this method, the mass spectra of triclopyr-TBDMS and 2-MP contain additional ions (Section H.2.) that may be used for confirmation.

3. Assay Time

A typical analytical run would consist of a minimum of four standards encompassing the expected range of sample concentrations, a reagent blank, a control (a non-fortified sample), a minimum of two fortified controls (one of which must be at the LOQ), and 15 samples. This typical analytical run could be prepared in approximately 24 hours (three working days).

There are several acceptable "stopping points" in the method, where sample preparation (Section I) may be suspended, upon completion of a step, without deleterious effects on the sample analysis. These are indicated below:

- a. Step I.1.p. In preparing large sets of samples, this is a convenient stopping point at the end of the first 8-hour day.
- b. Step I.1.q.
- c. Step I.1.r.(7).
- d. Step I.1 s.
- e. Step I.1.w. In preparing large sets of samples, this is a convenient stopping point at the end of the second 8-hour day.
- f. Step I.1.z.(8).
- g. Step I.1.gg
- h. Step I.1.mm

If the samples are to be stored overnight, the vials should be capped with PTFE-lined caps.

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4. Standardization of SPE Elution Profiles

Variation in the alumina and C₁₈ SPE columns may influence the elution profiles of triclopyr, 3,5,6-TCP, and 2-MP. It is necessary to obtain an elution profile for each lot of SPE columns used to ensure optimum recovery and clean-up efficiency. The following procedures can be used:

a. Alumina SPE Profile for 2-MP

- (1) To a 12-mL vial containing 5 mL of ethyl ether, add 50 μL of the 100-μg/mL spiking solution (Section G.1.d.).
- (2) Place an alumina SPE column on the vacuum manifold.
- (3) Rinse the SPE column with 5 mL of methanol. (Do not allow the column bed to dry.)
- (4) Condition the SPE column with 5 mL of ethyl ether. (Do not allow the column bed to dry.)
- (5) Transfer the sample solution from Step K.4.a.(1) to the SPE column and allow the sample to flow through the column by gravity alone, collecting 1-mL aliquots in 12-mL vials. (Do not allow the column bed to dry.)
- (6) Rinse the sample vial with 5 mL of ethyl ether, and when the sample solution in Step K.4.a.(5) is within 2 mm of the top of the column bed, transfer the rinse to the SPE column. Allow the solvent to flow through the column by gravity alone, and continue to collect 1-mL aliquots in 12-mL vials.
- (7) For each fraction collected, add 2.0 mL of 1-chlorobutane to the sample vial.
- (8) Concentrate the solutions to less than 0.8 mL (but not to dryness) using an N-Evap evaporator. (Notes L.2. and L.8.)
- (9) Proceed as described in Section I.1.u. through I.1.x.
- (10) Calculate the percent recovery for 2-MP as described in Section I.2.

A typical elution profile is illustrated in Figure 18. If the elution profile differs from that shown, adjust the volume of ethyl ether to be collected in Step I.1.r.(6).

b. Alumina SPE Profiles for Triclopyr and 3,5,6-TCP

- (1) To a 12-mL vial containing 5 mL of ethyl ether, add 10 μ L of the 100- μ g/mL spiking solution (Section G.1.d.).
- (2) Place an alumina SPE column on the vacuum manifold.
- (3) Rinse the SPE column with 5 mL of methanol. (Do not allow the column bed to dry.)
- (4) Condition the SPE column with 5 mL of ethyl ether. (Do not allow the column bed to dry.)

- (5) Transfer the sample solution from Step K.4.b.(1) to the SPE column and allow the sample to flow through the column by gravity alone, discarding the eluate. (Do not allow the column bed to dry.)
- (6) Rinse the sample vial with 2 mL of ethyl ether, and when the sample solution in Step K.4.b.(5) is within 2 mm of the top of the column bed, transfer the rinse to the SPE column. Allow the solvent to flow through the column by gravity alone, discarding the eluate.
- (7) Rinse the sample vial with 5 mL of acetone and transfer the rinse to the SPE column. Allow the solvent to flow through the column by gravity alone, discarding the eluate.
- (8) Dry the SPE column under vacuum for 5 minutes.
- (9) Elute the triclopyr and 3,5,6-TCP from the alumina SPE column by passing 50 mL of 0.1 N sodium hydroxide solution through the column at a flow rate of 1-2 mL/min, collecting 5.0-mL aliquots in 40-mL vials.
- (10) For each fraction collected, proceed as described in Section I.1.aa. through I.1.nn.
- (11) Calculate the percent recovery for triclopyr and 3,5,6-TCP as described in Section I.2.

Typical elution profiles are illustrated in Figure 18. If the elution profiles differ from those shown, adjust the volume of 0 1 N sodium hydroxide to be collected in Step I.1.r.(11).

- c C₁₈ SPE Profiles for Triclopyr and 3,5,6-TCP
 - (1) To two 40-mL vials containing 25 mL of 0.1 N hydrochloric acid, add 10 μL of the 100-μg/mL spiking solution (Section G.1.d.).
 - (2) Place two C₁₈ SPE columns on the vacuum manifold.
 - (3) Rinse each SPE column with 5 mL of acetonitrile. (Do not allow the column beds to dry.)
 - (4) Condition each SPE column with 5 mL of 0.1 N hydrochloric acid. (Do not allow the column beds to dry)
 - (5) Transfer the sample solutions from Step K.4.c.(1) to the SPE columns, and slowly pull the samples through the columns at a flow rate of 1-2 mL/min with the aid of vacuum. Discard the eluates without allowing the column beds to dry.
 - (6) Rinse the sample vials with 2 mL of 0.1 N hydrochloric acid, and transfer the rinses to the SPE columns. Slowly pull the rinse solutions through the columns at a flow rate of 1-2 mL/min with the aid of vacuum, and discard the cluates
 - (7) Dry each SPE column under vacuum for 1 minute.

(8) Elute the triclopyr and 3,5,6-TCP from one of the C₁₈ SPE columns with 10 mL of the 40% acetonitrile/59% water/1% 1.0 N hydrochloric acid solution, collecting 1.0-mL aliquots in 40-mL vials.

- Elute the triclopyr and 3,5,6-TCP from the other C₁₈ SPE column with 10 mL of the 80% acetonitrile/19% water/1% 1.0 N hydrochloric acid solution, collecting 1.0-mL aliquots in 40-mL vials.
- (9) For each fraction collected, proceed as described in Section I.1.aa. through I.1.nn.
- (10) Calculate the percent recovery for triclopyr and 3,5,6-TCP as described in Section I.2.

Typical elution profiles are illustrated in Figure 19. If the elution profiles differ from that shown, adjust the volume of the 40% acetonitrile/59% water/1% 1.0 N hydrochloric acid solution to be discarded in Step I.1.z.(5) or the volume of the 80% acetonitrile/19% water/1% 1.0 N hydrochloric acid solution to be collected in Step I.1.z.(8)

L. Notes

- Equipment, glassware, materials, reagents, and chemicals considered to be equivalent to
 those specified may be substituted with the understanding that their performance must
 be confirmed by appropriate tests. Common laboratory supplies are assumed to be
 readily available and are, therefore, not listed
- 2. The N-Evap evaporator should be set at a water bath temperature of 30 °C and a nitrogen flow rate of approximately 200 mL/min. At elevated water bath temperatures and/or flow rates, the 3,5,6-TCP and 2-MP may volatilize, thereby reducing recoveries.
- 3. The filters are used in the carrier gas supply lines to purify the helium entering the gas chromatograph.
- 4. While the sample preparation of the fish tissues for the determination of triclopyr and 3,5,6-TCP differs from that for the determination of 2-MP, the gas chromatographic/mass spectrometric conditions are the same.
- 5 Several tuning, or calibration, options are available for the Model 597X series of MSDs. The "Maximum Sensitivity Autotune" and "Mid-Mass Autotune" calibration features were found to consistently yield approximately 5-10 times the sensitivity compared to that of the "Standard Autotune" calibration. In addition, either of these calibrations should be followed with a "Usertune" calibration at m/z 131, 219, and 264.
- 6. The ions monitored for the analytes and their respective internal standards are grouped by retention time so that only those ions representative of a given compound are monitored at that compound's retention time

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- 7. Fish tissues containing bony material or shells will react with the 5.0 N hydrochloric acid and may necessitate further addition of 5.0 N hydrochloric acid to maintain a pH<1.
- 8. The measurement of a non-critical volume (one that does not affect the calculations) can be most easily accomplished by comparing the height (volume) in the sample vial with another vial containing the desired measured volume.
- In transferring the 1-chlorobutane layer, it is important not to remove any water.
 Contaminating the 1-chlorobutane with water will have deleterious effects on the derivatization and subsequent GC/MSD analysis

M. References

- 1. Freund, J. E.; Williams, F. J. Dictionary/Outline of Basic Statistics; Dover. New York, 1991; p 170.
- 2. Keith, L. H.; Crummett, W; Deegan, J, Jr.; Libby, R. A.; Taylor, J. K.; Wentler, G. Anal. Chem. 1983, 55, 2210-2218.

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Table I. Recovery of Triclopyr from Fish Tissues

Sample	Date of	Triclop	уг, μg/g	Percent	Statistical
Number	Analysis	Added	Found	Recovery	Calculations
SN17993002a	26-Jun-1997	0.000	0.000		
SN17993003b	3-Jul-1997	0.000	0.000		
SN18003302c	26-Jun-1997	0.000	0.000		
SN18003303d	3-Jul-1997	0.000	0 000		
SN20178201¢	26-Jun-1997	0.000	0.000		
SN20178202f	3-Jul-1997	0.000	0 000		
SN17993002	26-Jun-1997	0.003	< 0 010	NAg	,
SN17993003	3-Jul-1997	0.003	< 0 0 1 0	NA	
SN18003302	26-Jun-1997	0.003	< 0.010	NA	
SN18003303	3-Jul-1997	0.003	< 0 0 1 0	NA	
SN20178201	26-Jun-1997	0.003	< 0.010	NA	
SN20178202	3-Jul-1997	0.003	< 0.010	NA	
SN17993002	26-Jun-1997	0.010	0.0097	97	
SN17993002	26-Jun-1997	0.010	0.0094	94	
SN17993003	3-Jul-1997	0.010	0.0077	77	
SN17993003	3-Jul-1997	0.010	0.0092	92	
SN18003302	26-Jun-1997	0.010	0.0095	95	
SN18003302	26-Jun-1997	0.010	0.0097	97	
SN18003303	3-Jul-1997	0.010	0.0090	90	
SN18003303	3-Jul-1997	0.010	0.0095	95	$\bar{x} = 0.0095$
SN20178201	26-Jun-1997	0.010	0.0104	104	s = 0.0007
SN20178201	26-Jun-1997	0.010	0.0095	95	$(3s)^h = 0.0021$
SN20178202	3-Jul-1997	0.010	0.0102	102	$(10s)^{i} = 0.0071$
SN20178202	3-Jul-1997	0.010	0 0103	103	RSD = 7.5%
SN17993002	26-Jun-1997	0.050	0 0456	91	
SN17993003	3-Jul-1997	0.050	0.0465	93	
SN18003302	26-Jun-1997	0.050	0.0518	104	-
SN18003303	3-Jul-1997	0.050	0.0473	95	$\bar{x} = 0.0478$
SN20178201	26-Jun-1997	0.050	0.0466	93	s = 0.0022
SN20178202	3-Jul-1997	0.050	0.0488	98	RSD = 4.7%
SN17993002	26-Jun-1997	0.500	0.483	97	
SN17993003	3-Jul-1997	0.500	0 450	90	
SN18003302	26-Jun-1997	0.500	0 461	. 92	
SN18003303	3-Jul-1997	0.500	0.431	86	$\bar{x} = 0.451$
SN20178201	26-Jun-1997	0.500	0.436	87	s = 0.019
SN20178202	3-Jul-1997	0.500	0.445	89	RSD = 4.2%

Table I. (Cont.) Recovery of Triclopyr from Fish Tissues

Sample	Date of	Triclopyr, μg/g		F	Percent	Statistical
Number	Analysis	Added	Found	R	ecovery	Calculations
SN17993002	26-Jun-1997	5.00	4.562		91	
SN17993003	3-Jul-1997	5.00	4.614		92	
SN18003302	26-Jun-1997	5.00	4.266		85	
SN18003303	3-Jul-1997	5.00	4.539		91	$\bar{x} = 4489$
SN20178201	26-Jun-1997	5.00	4.318		86	s = 0.157
SN20178202	3-Jul-1997	5.00	4.633		93	RSD = 3.5%
				x =	93	
				s =	6	
			1	n =	30	

^a SN17993002—Bluegill edible tissue.

b SN17993003—Bluegill inedible tissue.

c SN18003302—Catfish edible tissue.

d SN18003303—Catfish inedible tissue.

c SN20178201—Crayfish edible tissue.

f SN20178202—Crayfish inedible tissue.

⁸ NA = not applicable. The residue was below the 0.010- μ g/g limit of quantitation.

h Calculated limit of detection.

¹ Calculated limit of quantitation.

Table II. Recovery of 3,5,6-Trichloro-2-pyridinol from Fish Tissues

Sample	Date of	3,5,6-T	CP, μg/g	Percent	Statistical
Number	Analysis	Added	Found	Recovery	Calculations
SN17993002a	26-Jun-1997	0.000	0.000		
SN17993003b	3-Jul-1997	0.000	0.000		
SN18003302°	26-Jun-1997	0.000	0.000		
SN18003303d	3-Jul-1997	0.000	0.000		
SN20178201c	26-Jun-1997	0.000	0.000		
SN20178202f	3-Jul-1997	0.000	0.000		
SN17993002	26-Jun-1997	0.003	< 0.010	NAg	
SN17993003	3-Jul-1997	0.003	< 0.010	NA	
SN18003302	26-Jun-1997	0.003	< 0.010	NA	
SN18003303	3-Jul-1997	0.003	< 0.010	NA	
SN20178201	26-Jun-1997	0.003	< 0.010	NA	
SN20178202	3-Jul-1997	0.003	< 0.010	NA	
SN17993002	26-Jun-1997	0.010	0.0092	92	
SN17993002	26-Jun-1997	0.010	0.0093	93	
SN17993003	3-Jul-1997	0.010	0.0115	115	
SN17993003	3-Jul-1997	0.010	0.0115	115	
SN18003302	26-Jun-1997	0.010	0.0094	94	
SN18003302	26-Jun-1997	0.010	0.0092	92	
SN18003303	3-Jul-1997	0.010	0.0100	100	
SN18003303	3-Jul-1997	0.010	0.0102	102	$\bar{x} = 0.0099$
SN20178201	26-Jun-1997	0.010	0.0104	104	s = 0.0009
SN20178201	26-Jun-1997	0.010	0.0102	102	$(3s)^h = 0.0026$
SN20178202	3-Jul-1997	0.010	0.0091	91	$(10s)^1 = 0.0086$
SN20178202	3-Jul-1997	0.010	0 0092	92	RSD = 8.6%
SN17993002	26-Jun-1997	0.050	0 0445	89	
SN17993003	3-Jul-1997	0.050	0 0463	93	
SN18003302	26-Jun-1997	0.050	0.0431	86	
SN18003303	3-Jul-1997	0.050	0.0451	90	$\bar{x} = 0.0450$
SN20178201	26-Jun-1997	0.050	0.0502	100	s = 0.0031
SN20178202	3-Jul-1997	0.050	0.0409	82	RSD = 7.0%
SN17993002	26-Jun-1997	0.500	0.406	81	
SN17993003	3-Jul-1997	0 500	0 422	84	
SN18003302	26-Jun-1997	0.500	0.379	76	
SN18003303	3-Jul-1997	0.500	0.409	82	$\ddot{x} = 0.412$
SN20178201	26-Jun-1997	0.500	0 473	95	s = 0.034
SN20178202	3-Jul-1997	0.500	0.383	77	RSD = 8.3%

Table II. (Cont.) Recovery of 3,5,6-Trichloro-2-pyridinol from Fish Tissues

Sample	Date of	3,5,6-TCP, µg/g		Percent	Statistical
Number	Analysis	Added	Found	Recovery	Calculations
SN17993002	26-Jun-1997	5.00	4.359	87	
SN17993003	3-Jul-1997	5.00	4.585	92	
SN18003302	26-Jun-1997	5.00	4.248	85	
SN18003303	3-Jul-1997	5.00	4.581	92	$\bar{x} = 4.484$
SN20178201	26-Jun-1997	5.00	4.911	98	s = 0.262
SN20178202	3-Jul-1997	5.00	4.220	84	RSD = 5.8%
			5	$\bar{x} = 92$	
				s = 9	
]	n = 30	

a SN17993002—Bluegill edible tissue.

b SN17993003—Bluegill inedible tissue

c SN18003302—Catfish edible tissue.

d SN18003303—Catfish inedible tissue.

[•] SN20178201—Crayfish edible tissue.

f SN20178202—Crayfish inedible tissue.

g NA = not applicable. The residue was below the $0.010-\mu g/g$ limit of quantitation.

h Calculated limit of detection.

i Calculated limit of quantitation

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Table III. Recovery of 2-Methoxy-3,5,6-trichloropyridine from Fish Tissues

Sample	Date of	2-MF	² , μg/g	Percent	Statistical
Number	Analysis	Added	Found	Recovery	Calculations
SN17993002 ^a	26-Jun-1997	0.000	0.000		
SN17993003b	3-Jul-1997	0.000	0.000		
SN18003302°	26-Jun-1997	0.000	0 003		
SN18003303d	3-Jul-1997	0.000	0.003		
SN20178201°	26-Jun-1997	0.000	0 000		
SN20178202f	3-Jul-1997	0.000	0.000		
SN17993002	26-Jun-1997	0.003	< 0 010	NAg	
SN17993003	3-Jul-1997	0.003	< 0.010	NA	
SN18003302	26-Jun-1997	0.003	< 0 010	NA	
SN18003303	3-Jul-1997	0.003	< 0.010	NA	
SN20178201	26-Jun-1997	0.003	< 0.010	NA	
SN20178202	3-Jul-1997	0.003	< 0.010	NA	
SN17993002	26-Jun-1997	0.010	0.0086	86	
SN17993002	26-Jun-1997	0.010	0.0089	89	
SN17993003	3-Jul-1997	0.010	0 0089	89	
SN17993003	3-Jul-1997	0.010	0.0095	95	
SN18003302	26-Jun-1997	0.010	0.0101	101	
SN18003302	26-Jun-1997	0.010	0.0098	98	
SN18003303	3-Jul-1997	0.010	0.0091	91	
SN18003303	3-Jul-1997	0.010	0 0091	91	$\bar{x} = 0.0093$
SN20178201	26-Jun-1997	0.010	0 0082	82	s = 0.0007
SN20178201	26-Jun-1997	0.010	0.0086	86	$(3s)^h = 0.0021$
SN20178202	3-Jul-1997	0.010	0.0098	98	$(10s)^{1} = 0.0069$
SN20178202	3-Jul-1997	0.010	0.0105	105	RSD = 7.5%
SN17993002	26-Jun-1997	0.050	0.0437	87	
SN17993003	3-Jul-1997	0.050	0.0439	88	
SN18003302	26-Jun-1997	0.050	0.0492	98	•
SN18003303	3-Jul-1997	0.050	0 0551	110	$\bar{x} = 0.0468$
SN20178201	26-Jun-1997	0 050	0 0430	86	s = 0.0046
SN20178202	3-Jul-1997	0.050	0 0458	92	RSD= 9.9%
SN17993002	26-Jun-1997	0.500	0.399	80	
SN17993003	3-Jul-1997	0.500	0.497	99	
SN18003302	26-Jun-1997	0.500	0.461	92	
SN18003303	3-Jul-1997	0.500	0 478	96	$\bar{x} = 0.442$
SN20178201	26-Jun-1997	0.500	0.383	77	s = 0.045
SN20178202	3-Jul-1997	0.500	0.432	86	RSD = 10.2%

Table III. (Cont.) Recovery of 2-Methoxy-3,5,6-trichloropyridine from Fish Tissues

Sample	Date of	2-MP, μg/g		Percen	t Statistical
Number	Analysis	Added	Found	Recove	ry Calculations
SN17993002	26-Jun-1997	5.00	4.103	82	
SN17993003	3-Jul-1997	5.00	5.066	101	
SN18003302	26-Jun-1997	5.00	4.580	92	
SN18003303	3-Jul-1997	5.00	4.968	99	$\bar{x} = 4.578$
SN20178201	26-Jun-1997	5.00	4.035	81	s = 0.431
SN20178202	3-Jul-1997	5.00	4.717	94	RSD = 9.4%
			<u></u>	$\bar{x} = 92$	
				s = s	
			1	n = 30	

a SN17993002—Bluegill edible tissue.

b SN17993003—Bluegill inedible tissue.

^c SN18003302—Catfish edible tissue. Recovery results from samples prepared using this sample were corrected for the amount of 2-MP found in the control.

d SN18003303—Catfish inedible tissue. Recovery results from samples prepared using this sample were corrected for the amount of 2-MP found in the control.

[•] SN20178201—Crayfish edible tissue.

f SN20178202—Crayfish inedible tissue.

^g NA = not applicable. The residue was below the $0.010-\mu g/g$ limit of quantitation.

h Calculated limit of detection

¹ Calculated limit of quantitation.

Triclopyr
Formula: C₇H₄Cl₃NO₃
Molecular Weight: 255

Triclopyr-TBDMS
Formula: C₁₃H₁₈Cl₃NO₃Si
Molecular Weight: 369

3,5,6-TCP Formula: C₅H₂Cl₃NO Molecular Weight: 197 3,5,6-TCP-TBDMS Formula: C₁₁H₁₆Cl₃NOSi Molecular Weight: 311

2-MP Formula: C₆H₄Cl₃NO Molecular Weight: 211

Figure 1. Chemical Structures of Triclopyr, 3,5,6-Trichloro-2-pyridinol and their TBDMS Derivatives, and 2-Methoxy-3,5,6-trichloropyridine

$$\begin{array}{c} \text{Cl} & \text{NH}_2 \\ \text{Cl} & \text{Cl} \\ \text{F} & \text{N} & \text{O} \end{array} \begin{array}{c} \text{OH} & \begin{array}{c} \text{MTBSTFA} \\ \text{60 °C, 1 hr.} \end{array} \end{array} \begin{array}{c} \text{Cl} \\ \text{F} & \text{N} & \text{O} \\ \text{O} & \begin{array}{c} \text{Cl} \\ \text{CH}_3 \end{array} \end{array}$$

Fluroxypyr Formula: C₉H₅Cl₂FN₂O₃ Molecular Weight: 254 Fluroxypyr-TBDMS Formula: C₁₃H₁₉Cl₂FN₂O₃Si Molecular Weight: 368

Fluroxypyr-DCP Formula: C₅H₃Cl₂FN₂O Molecular Weight: 196 Fluroxypyr-DCP-TBDMS Formula: C₁₁H₁₇Cl₂FN₂OSi Molecular Weight: 310

Fluroxypyr-MP Formula: C₆H₅Cl₂FN₂O Molecular Weight: 210

Figure 2. Chemical Structures of Fluroxypyr, 4-Amino-3,5-dichloro-6-fluoro-2-pyridinol and their TBDMS Derivatives, and 4-Amino-3,5-dichloro-6-fluoro-2-methoxypyridine

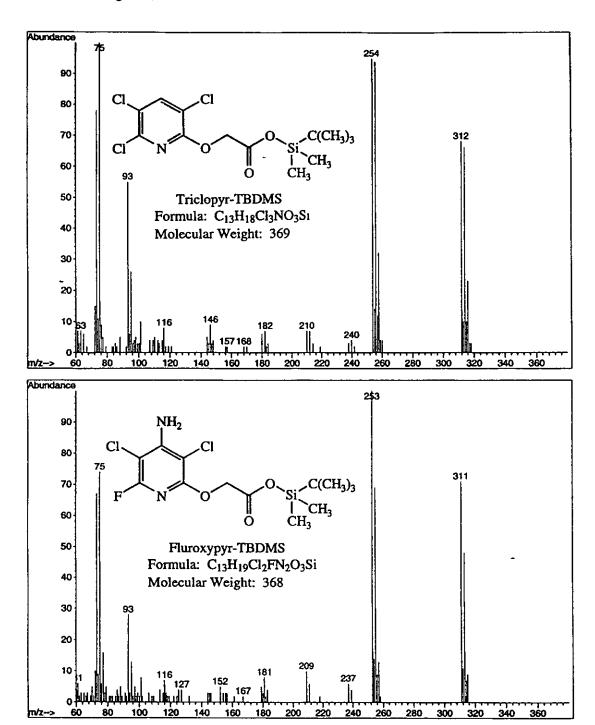


Figure 3. Mass Spectra of the TBDMS Derivatives of Triclopyr and Fluroxypyr

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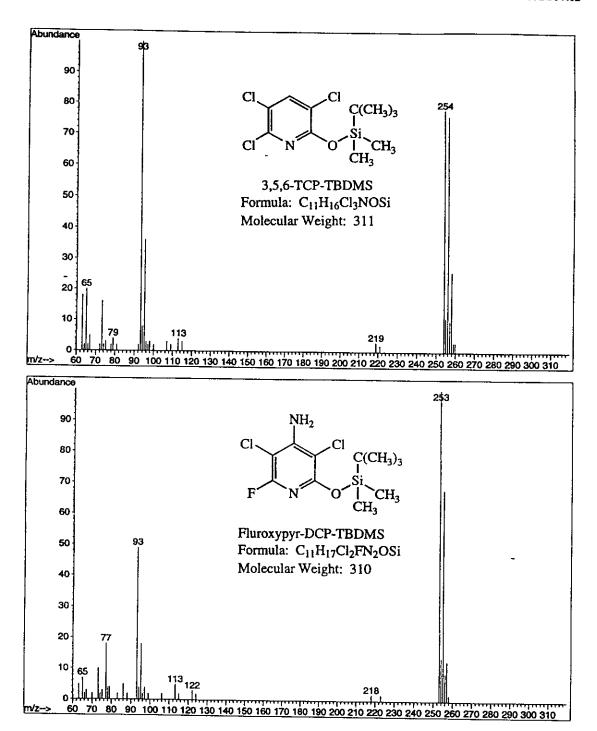


Figure 4. Mass Spectra of the TBDMS Derivatives of 3,5,6-Trichloro-2-pyridinol and 4-Amino-3,5-dichloro-6-fluoro-2-pyridinol

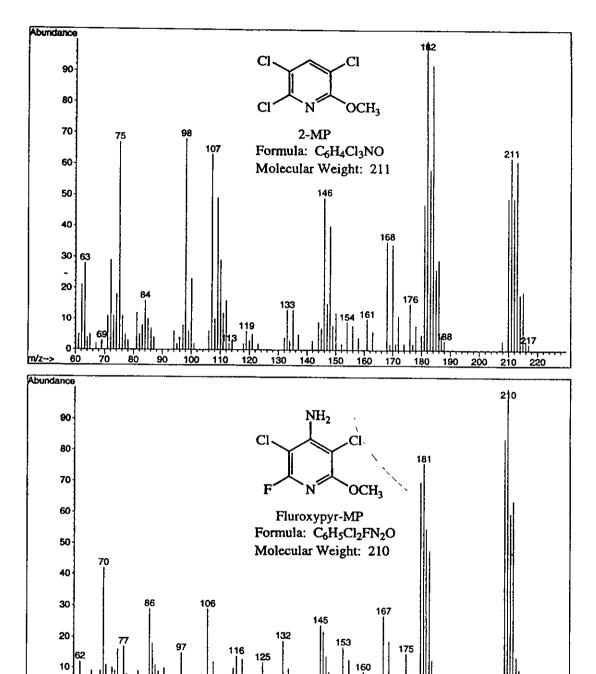


Figure 5. Mass Spectra of 2-Methoxy-3,5,6-trichloropyridine and 4-Amino-3,5-dichloro-6-fluoro-2-methoxypyridine

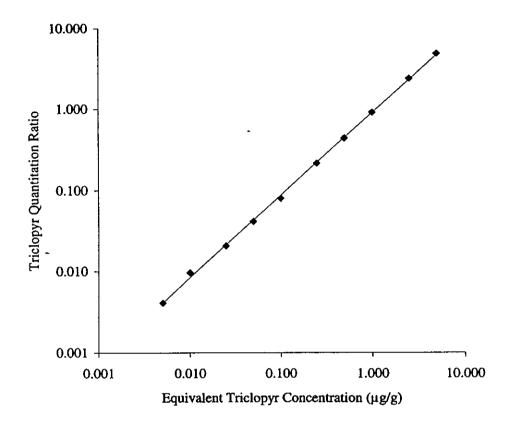
180 190

200

210

Effective Date: August 19, 1997

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Triclopyr Concentration	Equivalent Sample Conc.	Triclopyr Quantitation Ratio
μg/mL	μg/g	m/z 314 / m/z 311
0.005	0.005	0.00410
0.010	0.010	0.00961
0.025	0.025	0.02052
0.050	0.050	0.04120
0.100	0.100	0.07899
0.250	0.250	0.21348
0.500	0.500	0.43397
1.00	1.00	0.89599
2.50	2.50	2.33394
5.00	5.00	4.74820

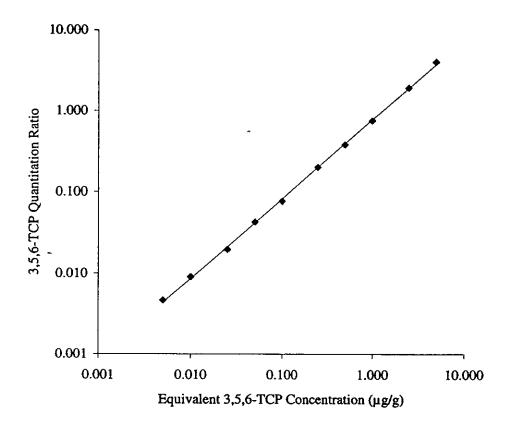
Power Regression Equation: $X = (Y/0.8925)^{(1/1.0139)}$

Correlation Coefficient (r²): 0.99934

Figure 6. Typical Calibration Curve for the Determination of Triclopyr in Fish Tissues

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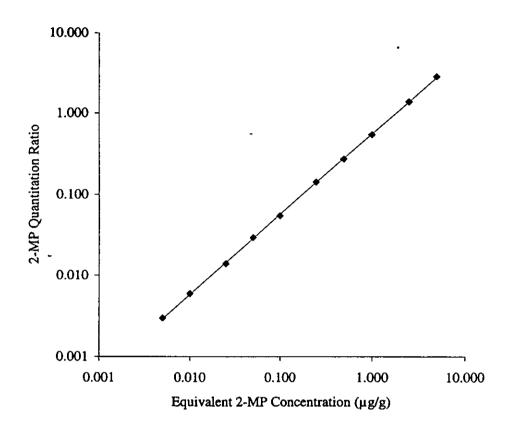


3,5,6-TCP Concentration	Equivalent Sample Conc.	3,5,6-TCP Quantitation Ratio
μg/mL	μg/g	m/z 254 / m/z 253
0.005	0.005	0.00466
0.010	0.010	0.00904
0.025	0.025	0.01964
0.050	0.050	0.04240
0.100	0.100	0.07636
0.250	0.250	0.19917
0.500	0.500	0.38024
1.00	1.00	0.74854
2.50	2.50	1.90305
5.00	5.00	3.99394

Power Regression Equation: $X = (Y/0.7720)^(1/0.9755)$

Correlation Coefficient (r²): 0.99950

Figure 7. Typical Calibration Curve for the Determination of 3,5,6-Trichloro-2-pyridinol in Fish Tissues

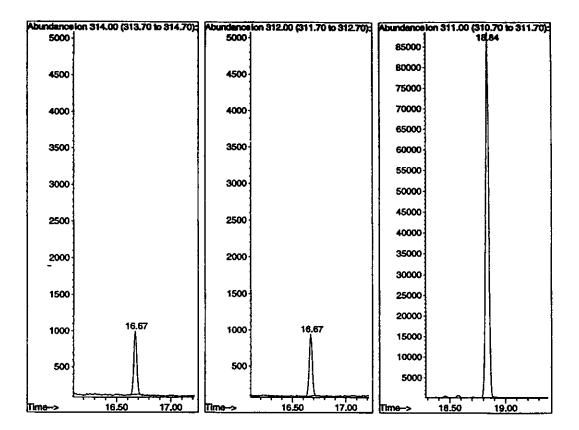


2-MP Concentration µg/mL	Equivalent Sample Conc. µg/g	2-MP Quantitation Ratio m/z 213 / m/z 210
0.005	0.005	0.00298
0.010	0.010	0.00594
0.025	0.025	0.01391
0.050	0.050	0.02918
0.100	0.100	0.05408
0.250	0.250	0.14065
0.500	0.500	0.27228
1.00	1.00	0.54151
2.50	2.50	1.37561
5.00	5 00	2.77943

Power Regression Equation: $X = (Y/0.5504)^{(1/0.9886)}$

Correlation Coefficient (r²): 0.99988

Figure 8. Typical Calibration Curve for the Determination of 2-Methoxy-3,5,6-trichloropyridine in Fish Tissues



Data File : 0601006.D

ALS Bottle : 6

Date : 3 Jul 97 18:27
Data Path : C:\HPCHEMPC\1\DATA\RES94084\A070397A.ELO\

Instrument : GC/MSD S/N 3307A00401

Protocol : RES94084

Sample Name: Triclopyr Standard - 0010 ng/mL Sample Info: Equivalent to 0010 ng/g in fish

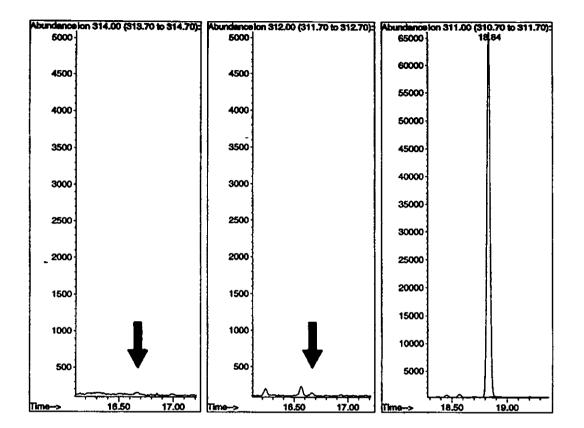
Operator : Edward L. Olberding

INTERNAL STANDARD RETENTION TIME: 18.84 PEAK AREA (M/Z 311) : 1723923

TRICLOPYR RETENTION TIME 16.67 PEAK AREA (M/Z 314) 16562 PEAR AREA (M/Z 312) 16597

Equivalent Triclopyr Concentration: 0.0100 μg/g

Figure 9. Typical Chromatogram of a 0.010-µg/mL Standard Equivalent to 0.010 µg/g of Triclopyr in Fish Tissue



Data File : 1901019.D

ALS Bottle : 19

Date : 4 Jul 97 1:05

Data Path : C:\HPCHEMPC\1\DATA\RES94084\A070397A.ELO\

Instrument : GC/MSD S/N 3307A00401

Protocol : RES94084

Sample Name: SN20178202 - Control Sample Info: Sample 26 - Crayfish - Inedible

Operator : Edward L. Olberding

INTERNAL STANDARD RETENTION TIME: 18.84 : 1323914 PEAK AREA (M/Z 311)

NO TRICLOPYR FOUND

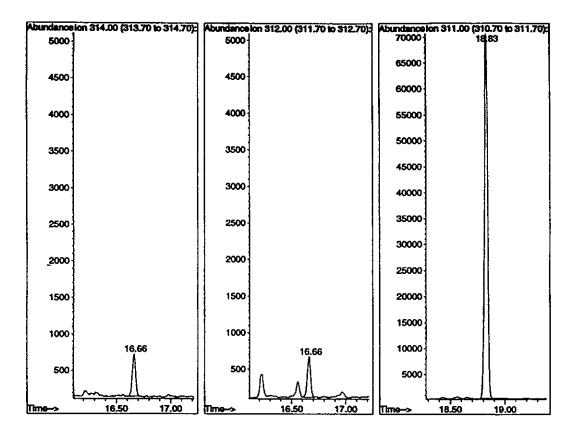
Triclopyr Concentration: 0.0000 µg/g

Figure 10. Typical Chromatogram of a Control Crayfish Inedible Tissue Sample Containing No Detectable Residue of Triclopyr

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Data File : 2901029.D

ALS Bottle : 29

Date : 4 Jul 97 6:10
Data Path : C:\HPCHEMPC\1\DATA\RBS94084\A070397A.ELO\

Instrument : GC/MSD S/N 3307A00401

Protocol : RES94084

Sample Name: SN20178202 - Spiked at 0010 ng/g Sample Info: Sample 34 - Crayfish - Inedible

Operator : Edward L. Olberding

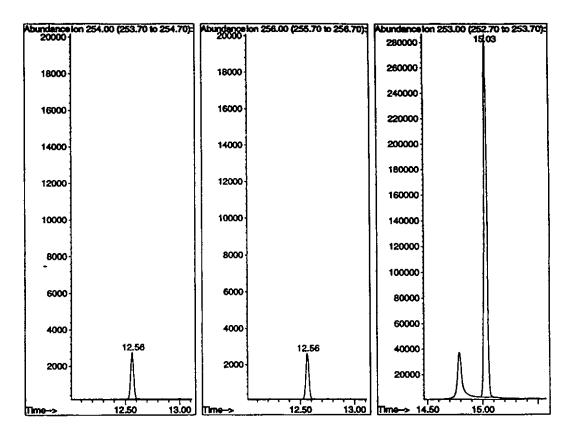
INTERNAL STANDARD RETENTION TIME: 18.83 PEAK AREA (M/Z 311) 1443236 TRICLOPYR RETENTION TIME 16.66 PEAK AREA (M/Z 314) 12260 PEAK AREA (M/Z 312) 11953

Triclopyr Concentration: 0.0102 µg/g

Recovery: 102%

Figure 11. Typical Chromatogram of a Crayfish Inedible Tissue Sample Fortified with 0.010 µg/g of Triclopyr

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Data File : 0601006.D

ALS Bottle : 6

: 3 Jul 97 18:27 Date

Data Path : C:\HPCHEMPC\1\DATA\RES94084\A070397A.ELO\
Instrument : GC/MSD S/N 3307A00401

: RES94084 Protocol

Sample Name: Triclopyr Standard - 0010 ng/mL Sample Info: Equivalent to 0010 ng/g in fish

: Edward L. Olberding Operator |

INTERNAL STANDARD RETENTION TIME: 15.03 PEAK AREA (M/Z 253)

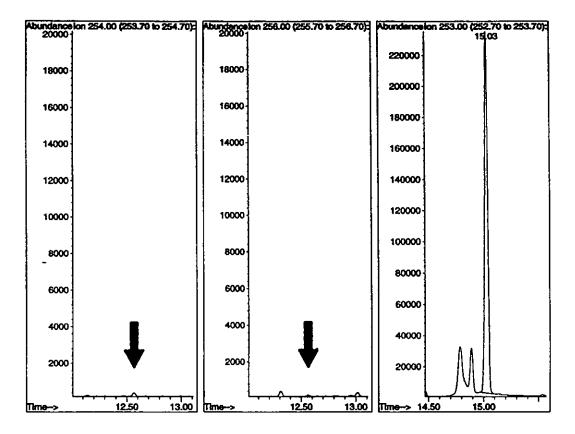
3,5,6-TCP RETENTION TIME 12.56 : PEAK AREA (M/Z 254) 49230 PEAK AREA (M/Z 256) 48808

Equivalent 3,5,6-TCP Concentration: 0.0100 μg/g

Figure 12. Typical Chromatogram of a 0.010-μg/mL Standard Equivalent to 0.010 μg/g of 3.5.6-TCP in Fish Tissue

Effective Date: August 19, 1997

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Data File : 1901019.D

ALS Bottle : 19

4 Jul 97 1:05 Date

Date : 4 Jul 97 1:05
Data Path : C:\HPCHEMPC\1\DATA\RES94084\A070397A.ELO\

Instrument : GC/MSD S/N 3307A00401

Protocol : RES94084

Sample Name: SN20178202 - Control Sample Info: Sample 26 - Crayfish - Inedible

Operator : Edward L. Olberding

INTERNAL STANDARD RETENTION TIME: 15.03 : 4486369 PEAK AREA (M/Z 253)

NO 3,5,6-TCP FOUND

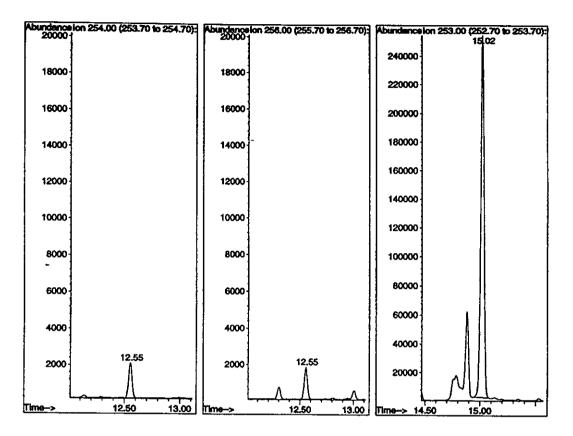
3,5,6-TCP Concentration: 0.0000 µg/g

Figure 13. Typical Chromatogram of a Control Crayfish Inedible Tissue Sample Containing No Detectable Residue of 3.5.6-TCP

DowElanco Study ID: RES94084

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Data File : 2901029.D

ALS Bottle : 29

Date : 4 Jul 97 6:10
Data Path : C:\HPCHEMPC\1\DATA\RES94084\A070397A.ELO\

Instrument : GC/MSD S/N 3307A00401

Protocol : RES94084

Sample Name: SN20178202 - Spiked at 0010 ng/g Sample Info: Sample 34 - Crayfish - Inedible

Operator : Edward L. Olberding

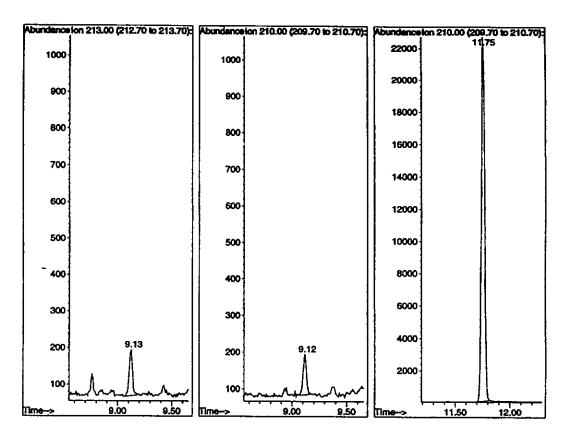
INTERNAL STANDARD RETENTION TIME: 15.02 PEAK AREA (M/Z 253) 5192065

3,5,6-TCP RETENTION TIME 12.55 PEAK AREA (M/Z 254) PEAK AREA (M/Z 256) 41066 36079

3,5,6-TCP Concentration: 0.0091 µg/g

Recovery: 91%

Figure 14. Typical Chromatogram of a Control Crayfish Inedible Tissue Sample Fortified with $0.010 \,\mu g/g$ of 3,5,6-TCP



Data File : 0601006.D

ALS Bottle : 6

Date : 5 Jul 97 12:20
Data Path : C:\HPCHEMPC\1\DATA\RES94084\A070597A.ELO\

Instrument : GC/MSD S/N 3307A00401

Protocol : RES94084

Sample Name: Triclopyr Standard - 0010 ng/mL Sample Info: Equivalent to 0010 ng/g in fish

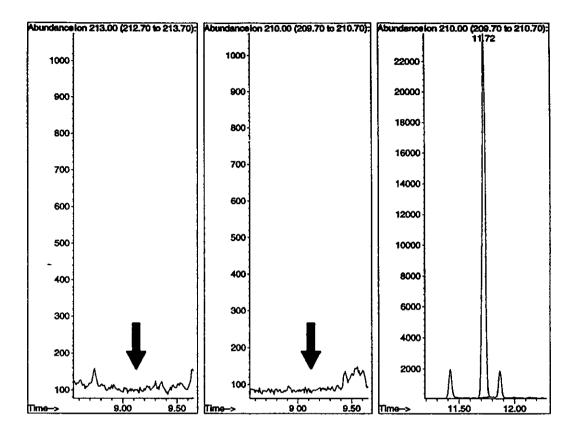
Operator : Edward L. Olberding

INTERNAL STANDARD RETENTION TIME: 11.75 PEAK AREA (M/Z 210) : 465618

2-MP RETENTION TIME 9.13 PEAK AREA (M/Z 213) 2765 PEAK AREA (M/Z 210) 2357

Equivalent 2-MP Concentration: 0.0100 μg/g

Figure 15. Typical Chromatogram of a 0.010-μg/mL Standard Equivalent to 0.010 μg/g of 2-MP in Fish Tissue



Data File : 1901019.D ALS Bottle : 19 Date : 5 Jul 97 18:04

Data Path : C:\HPCHEMPC\1\DATA\RES94084\A070597A.ELO\

Instrument : GC/MSD S/N 3307A00401

Protocol : RES94084

Sample Name: SN20178202 - Control Sample Info: Sample 26 - Crayfish - Inedible

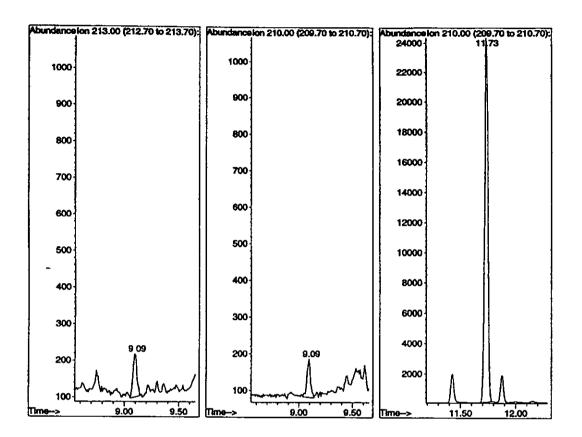
Operator : Edward L. Olberding

INTERNAL STANDARD RETENTION TIME: PEAK AREA (M/Z 210) :

NO 2-MP FOUND

2-MP Concentration: 0.0000 µg/g

Figure 16. Typical Chromatogram of a Control Crayfish Inedible Tissue Containing No Detectable Residue of 2-MP



Data File : 3001030.D

ALS Bottle : 30

Date : 5 Jul 97 22:56
Data Path : C:\HPCHEMPC\1\DATA\RES94084\A070597A.ELO\

Instrument : GC/MSD S/N 3307A00401

: RES94084 Protocol

Sample Name: SN20178202 - Spiked at 0010 ng/g Sample Info: Sample 35 - Crayfish - Inedible

Operator : Edward L. Olberding

INTERNAL STANDARD RETENTION TIME: 11.73 PEAK AREA (M/Z 210) 492706

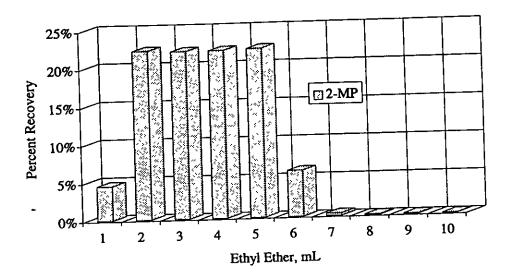
2-MP RETENTION TIME 9.09 PEAK AREA (M/Z 213) PEAK AREA (M/Z 210) 3002 2444

2-MP Concentration: 0.0102 µg/g

Recovery: 102%

Figure 17. Typical Chromatogram of a Control Crayfish Inedible Tissue Fortified with 0.010 µg/g of 2-MP

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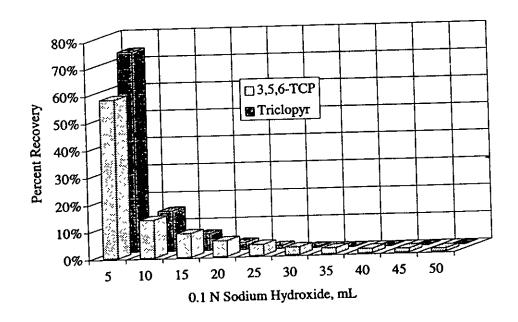
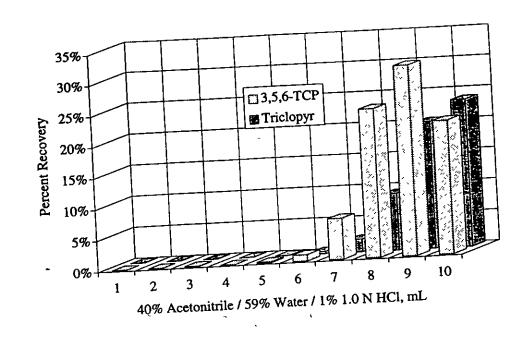


Figure 18. Typical Alumina SPE Elution Profiles for 2-Methoxy-3,5,6-trichloropyridine (Ethyl Ether) and Triclopyr and 3,5,6-Trichloro-2-pyridinol (0.1 N Sodium Hydroxide)

GRM 97.02 Effective Date: August 19, 1997



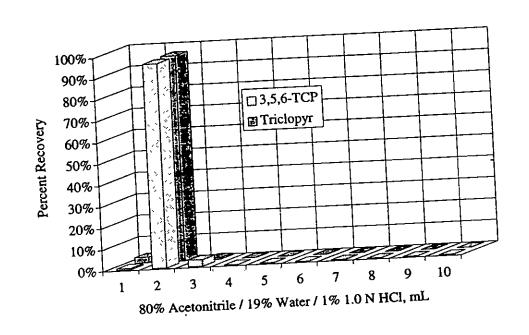


Figure 19. Typical C₁₈ SPE Elution Profiles for Triclopyr and 3,5,6-Trichloro-2-pyridinol

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Conditions and Substitutions used by ACB

- 1. The ACB used the following instrumentation: HP model 5890 Series II plus gas chromatograph, HP model 5972 Mass Selective Detector ACB used a Restek Rtx-1701 fused silica capillary column, 10 m x 0.18 mm i.d , 0.4 μ m film thickness (Cat. # 42010). The temperature program was the same as the petitioner: 70 °C for 1.0 min, 70 °C to 255 °C at 10 °C/min, 255 °C to 280 °C at 20 °C/min, and 280 °C for 4.75 min. The injector and interface temperatures were 280 °C The carrier gas was helium. The injection volume was 1 μ L.
- 2. The ACB purchased graduated 10 mL Reacti-Vials™ from Pierce Chemical Co, Rockford, Illinois 61105 (Cat.#13225) for the final extraction step. This substitution eliminated the need to estimate the 1 mL final volume in the 12 mL unmarked vial used in the method.
- 3a. The method stated to profile each lot of SPE columns prior to use ACB found only one lot of Supelco, Inc. (the method calls for Bakerbond spe TM) C_{18} columns that would give adequate recoveries to use in the trial. The method addresses the use of equivalent reagents with the understanding that they must be confirmed by appropriate tests. The ACB contacted the petitioner to discuss the low recovers from the profiling results. The petitioner suggested the use of other manufacturers of C_{18} columns. In addition, the petitioner gave the ACB permission to increase the percentage of acetonitrile (ACN) in the elution solution to increase recoveries. The ACB used an elution solution: ACN-H₂O-1N HCI (90+9+1, v/v/v). The method used ACN-H₂O-1N HCI (80+19+1, v/v/v). The ACB suggests that a column profile be performed using a standard that mirrors the tolerance level as well as the level (1µg/mL) in the profiling instructions.
- 3b. The ACB found that 5 mL of elution solution was a more efficient volume than the 4 mL used in the method (step I.1.y.(8)). In addition, the ACB collected a second elution volume (5 mL) in another 40 mL vial and stored it capped in the refrigerator. This could be worked up later if the recoveries were low (<70%) for the first aliquot. This is a recommendation for higher tolerance levels
- 4 The method did not give storage or stability data for the standard solutions or the derivatized calibration standards. The ACB stored stock standards and derivatized calibration, standards in the refrigerator at 2-11°C. The ACB found that the final extracts were stable at room temperature for a least 48 hours in capped auto-sampler vials.
- 5. The method emphasizes, in a note (L.9), that it is critical not to-transfer any water over with the 1-chlorobutane layer because it will have a deleterious effect on the derivatization and subsequent GC/MSD analysis. The ACB observed as much as a ten fold difference in area responses when aqueous (salt) was in the final extract.

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6 The method suggested the use of a calibration exponential regression equation to determine sample concentrations (quantitation ratio vs concentration). The ACB ran calibration curves to demonstrate linearity over the dynamic range of the fortified samples. The ACB used the standard regression equation (Y = mx + b). In this method validation, The ACB determined sample concentrations from a ratio of sample responses to the average of standard responses that bracketed the samples during the analysis